





CC MOLECULE TYPE: DNA (genomic)  
S0 SEQUENCE 965 BP; 192 A; 170 C; 226 G; 200 T; 177 OTHER.  
Query Match 6.2%; Score 34; DB 3; Length 965;  
Best Local Similarity 12.7%; Pred. No. 2.66e-08;  
Matches 20; Conservative 78; Mismatches 58; Indels 2; Gaps 2;  
Db 777 GTKGRUHVHVGSGVRSSTCTASDYYTSTYWGVRGKMGWDYGGGYNYNKGKRVTVMA 836  
CP 7386 GTTGGAGGAGGAGGACATGCAATCAGACTGGAAGAGTATCAACAAGTTGGTTCAA 7327  
Db 837 DTSSNSRBS-VTAADTAYYYCVGRSYDSGDYWGCTTVYSSHUVYDMSS-SSASVG 894  
CP 7326 CCCGGGGCTTGTAATCCAGAGGGCGAGAGAGGCCAAGGCTGCACCTTGAGGAGAC 7267  
Db 895 DRVYTCRSSTTHGNGNTYYWYKAKARYVSNRSVRS 932  
CP 7266 TGCAGAGGGCAAGCCCAACAGGTTCTCTCTACAG 7229  
RESULT 4  
ID US-08-388-672A-22 STANDARD; DNA; UNC; 965 BP.  
AC xxxxxx  
DE Sequence 22, Application US/08388672A  
CC Sequence 22, Application US/08388672A  
CC Patent No. 5795961  
CC GENERAL INFORMATION:  
CC APPLICANT: Wallace, T. Paul  
CC APPLICANT: Harliss, William J.  
CC APPLICANT: Carr, Frank J.  
CC APPLICANT: Old, Lloyd J.  
CC APPLICANT: Welt, Sydney  
CC APPLICANT: Kitamura, Kunio  
CC TITLE OF INVENTION: Recombinant Human Anti-Lewis B  
CC NUMBER OF SEQUENCES: 25  
CC CORRESPONDENCE ADDRESS:  
CC ADDRESSEE: Felife and Lynch  
CC STREET: 805 Third Avenue  
CC CITY: New York  
CC STATE: New York  
CC COUNTRY: U.S.A.  
CC ZIP: 10022  
CC COMPUTER READABLE FORM:  
CC MEDIUM TYPE: Floppy disk  
CC COMPUTER: IBM PC compatible  
CC OPERATING SYSTEM: PC-DOS/MS-DOS  
CC SOFTWARE: Patentin Release #1.0, Version #1.30  
CC CURRENT APPLICATION DATA:  
CC APPLICATION NUMBER: US/08/388,672A  
CC FILING DATE: 14-FEB-1995  
CC CLASSIFICATION:  
CC ATTORNEY/AGENT INFORMATION:  
CC NAME: Hanson, NO. 5795961man D.  
CC REGISTRATION NUMBER: 30,946  
CC REFERENCE/DOCKET NUMBER: LUD 5409  
CC TELECOMMUNICATION INFORMATION:  
CC TELEPHONE: 212-688-9200  
CC TELEFAX: 212-838-3884  
CC INFORMATION FOR SEQ ID NO: 22:  
CC SEQUENCE CHARACTERISTICS:  
CC LENGTH: 965 base pairs  
CC TYPE: nucleic acid  
CC STRANDEDNESS: unknown  
CC TOPOLOGY: unknown  
CC MOLECULE TYPE: DNA (genomic)  
S0 SEQUENCE 965 BP; 192 A; 170 C; 226 G; 200 T; 177 OTHER.  
Query Match 5.8%; Score 32; DB 3; Length 965;  
Best Local Similarity 13.2%; Pred. No. 4.83e-07;  
Matches 15; Conservative 56; Mismatches 43; Indels 0; Gaps 0;

Db 837 DTSSNSRBSVTAADTAYYYCVGRSYDSGDYWGCTTVYSSHUVYDMSSSSASVDR 896  
OY 7387 ATGCACTGCAATATGAGACCACCTGCAGATGCTGCCTCAGCCAAAGAGCTGGGAGGA 7446  
Db 897 VTTCCSSTTHGNGNTYYWYKAKARYVSNRSVRSRSGSGNDYTTSDATYYC 950  
OY 7447 CTGTGACAGGACTGCAGGACGAGGGGACAGGGTTGTCTCTCGGTACTTC 7500  
RESULT 5  
ID US-08-232-463-14 STANDARD; DNA; UNC; 7218 BP.  
AC xxxxxx  
DE Sequence 14, Application US/08232463  
CC Sequence 14, Application US/08232463  
CC Patent No. 5670367  
CC GENERAL INFORMATION:  
CC APPLICANT: DORNER, F.  
CC APPLICANT: SCHEFLINGER, F.  
CC APPLICANT: FALKNER, F. G.  
CC TITLE OF INVENTION: RECOMBINANT FOWLPOX VIRUS  
CC NUMBER OF SEQUENCES: 52  
CC CORRESPONDENCE ADDRESS:  
CC ADDRESSEE: Foley & Lardner  
CC STREET: 1800 Diagonal Road, Suite 500  
CC CITY: Alexandria  
CC STATE: VA  
CC COUNTRY: USA  
CC ZIP: 22313-0299  
CC COMPUTER READABLE FORM:  
CC MEDIUM TYPE: Floppy disk  
CC COMPUTER: IBM PC compatible  
CC OPERATING SYSTEM: PC-DOS/MS-DOS  
CC SOFTWARE: Patentin Release #1.0, Version #1.25  
CC CURRENT APPLICATION DATA:  
CC APPLICATION NUMBER: US/08/232,463  
CC FILING DATE:  
CC CLASSIFICATION: 435  
CC PRIOR APPLICATION DATA:  
CC APPLICATION NUMBER: US/07/935,313  
CC FILING DATE:  
CC APPLICATION NUMBER: EP 91 114 300.6  
CC FILING DATE: 26-AUG-1991  
CC ATTORNEY/AGENT INFORMATION:  
CC NAME: BENT, Stephen A.  
CC REGISTRATION NUMBER: 29,768  
CC REFERENCE/DOCKET NUMBER: 30472/114 IMMU  
CC TELECOMMUNICATION INFORMATION:  
CC TELEPHONE: (703)683-4109  
CC TELEFAX: (703)836-9300  
CC TELEX: 899149  
CC INFORMATION FOR SEQ ID NO: 14:  
CC SEQUENCE CHARACTERISTICS:  
CC LENGTH: 7218 base pairs  
CC TYPE: nucleic acid  
CC STRANDEDNESS: single  
CC TOPOLOGY: linear  
CC IMMEDIATE SOURCE:  
CC CLONE: pTZgpt-F15  
S0 SEQUENCE 7218 BP; 1944 A; 1491 C; 1486 G; 1929 T; 368 OTHER.  
Query Match 5.8%; Score 32; DB 2; Length 7218;  
Best Local Similarity 2.0%; Pred. No. 4.83e-07;  
Matches 2; Conservative 63; Mismatches 33; Indels 0; Gaps 0;  
Db 1339 YY 1398  
CP 7482 CCGTGTCCCTCCGCTCCGCTGAGTCTCCTGCCACAGTCCACACCTCTGGCTGAGG 7423  
Db 1399 YY 1436  
CP 7422 CCAGCATCTGCAGGTGGGTCTCATTTTCAGTCCATGT 7385









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SQ SEQUENCE 82 BP; 1 A; 2 C; 10 G; 8 T; 61 OTHER.

Query Match 4.58; Score 25; DB 4; Length 82;

Best Local Similarity	1.0%	Freq. NO.	1.13E-03;		
Matches	5; Conservative	20; Mismatches	41; Indels	0; Gaps	0;

Db 10 GAGNNBNNBNNBNNBNNBNNBNNBNNBNNBNNBNN 69

CP 7744 GAGACCCCTCTGGGGGAGAGGGGCTCTGGATCACCCTACTTCTCGCTGTCAGTTA 7685

Db 70 BNNBG 75

CP 7684 GGGGGG 7679

Search completed: Fri Jan 14 18:12:58 2000  
Job time : 93 secs.

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Job time : 93 secs.